



TABLE 5 (SEQ ID No:27 and SEQ ID No:28)

1 GGGGTGCANA GAAAGACAC CAGGCCCCAG CTGGAGGTG CTAACTCCAG AGGCCAGCAT CAGCAACTGG GCACAGAAAG

81 GAGCCGCTTG GGCAGGGACC ¹ATGGCAGGC CACATCCCTG GTGGCTGTGC GTTCTGGGA CCTGTGGG GCTCTACGT
M A R P H P W L C V L G T L V G L S A

161 ACTCCAGCCC CCAAGAGCTG CCCAGAGAG CACTACTGGG CTCAGGAAA GCTGTGCTGC CAGATGTGTG AGCCAGAAC
T P A P K S C P E R H Y W ~~W~~A Q G K L C C Q M C E P G T

241 ATTCCTCGTG AAGGACTGTG ACCAGCATAG AAAGGCTGCT CAGTGTGATC CTTCATACC GGGGCTCTCC TTCTCTCCTG
F L V K D C D Q H R K A A Q C D P C I P G V S F S P D

321 ACCACCAAC CCGCCCCAC TGTAGAGCT GTGGCACTG TAATCTGCT CTTCCTGTTG GCAACTGCAC CATCACTGCC
H H T R P H C E S C R H C N S G L L V R N C T I T A

401 AATGCTGAGT GTGCTGTG CAACTGGCTGG CAGTGCAGG ACAAGAGTG CACCGAGTGT GATCCTCTTC CAAACCTTC
N A E C A C R N G W Q C R D K E C T E C D P L P N P S

481 GCTGACCGCT CGTGCTCTC AGGCCCTGAG CCCACACCCT CAGCCCCACC ACTTACCTTA TGTAGTAG ATGCTGAGG
L T A R S S Q A L S P H P Q P T H L P Y V S E M L E A

561 CCAGGACAGC TGGGCACATG CAGACTCTGG CTGAQTTCAG GCAGCTGCCT GCCCGGACTC TCTCTACCA CTGGCCACC
R T A G H M Q T L A D F R Q L P A R T L S T H W P P

641 CAAGATCCC TGTGACGCTC CGATTTTATT CGCATCCTTG TGAATCTCTC TGAATGTTT CTGTGTTTCA CCCTGGCCGG
Q R S L C S S D F I R I L V I F S G M F L V F T L A G
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721 GGGCCTGTTT CTCCATCAAC GAAGAAATA TAGATCAAAC AAAGAGAAA GTCTGTGGA GCCTGCAGAG CCTTGTGTT
A L F L H Q R R K Y R S N K G E S P V E P A E P C R Y
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801 ACAGCTGCCC CAGGGAGGAG GAGGGACGA CCATCCCAT CCAGAGAGAT TACCGAAAC CGGAGCCTGC CTGCTCCCC
S C P R E E E G S T I P I Q E D Y R K P E P A C S P

881 TGAGCCAGCA CCTGCGGTAG CTGCACTACA GGCCTGGCCT CAACCCCAAC CCGCCGAC ATCCAAGGA GAGTAGAAC
* TGAGCCAGCA AACTGCACTC CCATCCTCTT GTCAAGGGCC TTTCCTGTGT ACACTGACA GAGTGCCCTTT TCGAGACTGG

1041 CAGGGACGAG GACAAATATG GATGAGGTGG AGAGTGGGA GCAAGAGCCC AGCCAAGCTGC CCGCGCCTGC AGGAGGGCGG

1121 GGGCTCTGCT TGTAAAGCAC ACTTCTGCT GCGAAGACC CACATGCTAC AAGACGGCA AATTAAGTG ACAGATGACC